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(54) Title: ATTENUATED VIRUSES AND METHOD OF MAKING THE SAME

(57) Abstract

Disclosed are attenuated viruses, not naturally occurring, that contain one or more additional methylation sites in the genome of the virus compared to the corresponding wild-type virus. Preferably, the methylation sites are added into the genome of the virus by introducing an additional CG segment into the genome by means of a silent mutation. The attenuated viruses are useful for producing an immune response, including both the production of antibodies in animals for diagnostic use and the induction of protective immunity in a subject. Pharmaceutical formulations and methods of making the attenuated viruses are also disclosed.

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ATTENUATED VIRUSES AND METHOD OF MAKING THE SAME

This invention was made with Government support under Grant No. RO1 CA47217 from the National Cancer Institute. The Government has certain rights to this invention.

Field of the Invention

This application concerns singly or multiply attenuated viruses useful as vaccines, where one applied attenuation strategy is to create additional sites for DNA methylation in the viral genome, such additional sites (1) not affecting the amino acid sequence of the virus, and (2) conferring improved host cell control over the expression of the viral genome.

Background of the Invention

There are at present no vaccines available

which are effective against human retroviral infections,
and only one which is effective in animals (feline
leukemia virus). Various strategies are currently being
investigated in attempts to develop effective vaccines
against viruses such as the human (HIV) and simian (SIV)

immunodeficiency viruses, including subunit vaccines and
whole or partial virus vaccines. Clinical trials of
potential HIV-1 vaccines have produced almost universal
failure; over a dozen large projects, utilizing either
peptide vaccines (small fragments of HIV-1 protein,
usually the glycoprotein coat) or killed, denatured
virus, have failed.

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Studies in non-human primates have demonstrated that removal of the nef gene from SIV immunizes monkeys against secondary challenge to SIV. A natural experiment appears to have likewise verified that removal of the nef gene may produce an effective live attenuated vaccine in humans. A population of individuals have been identified who have been infected with HIV for more than one decade, but who show no signs of progressing to AIDS. When the virus infecting these people was isolated and sequenced, it was determined that these particular HIV strains were spontaneous mutations at the nef gene locus; that is, the nef gene had undergone spontaneous deletion. Herein we describe a method to attenuate viruses used in vaccines whose genomes are a target for host cell DNA methylation.

Summary of the Invention

The present invention is based on the discovery DNA methylation sites, in contrast to other dinucleotides, have been preferentially lost during HIV-1 evolution at a rate which far surpasses that of host 20 genes. There is also a loss of methylation sites in DNA DNA methylation is a viruses and some RNA viruses. process by which the five position carbon atom of specific cytosines in DNA are methylated to create 5methylcytosine. In animal cells, most methylation occurs 25 in the CpG dinucleotide; that is, in cytosines which are immediately 5' to guanines. Generally, when genes are methylated, they are transcriptionally "silent" -- no messenger RNA and hence no protein is produced from them. The present invention employs the active introduction of 30 silent mutations (i.e., that do not affect the amino acid sequence) into the virus genome, such mutations creating methylation sites not normally present, methylation of which will impede viral function.

Accordingly, a first aspect of the present invention is an attenuated virus (or "modified virus"), not naturally occurring, containing at least 1 additional

methylation site introduced by mutation in the genome of the virus over the corresponding wild-type virus.

A second aspect of the present invention is a DNA encoding a virus as given above (e.g., a cDNA encoding a virus), as well as a vector (e.g., an expression vector) containing the DNA.

A third aspect of the present invention is a pharmaceutical formulation comprising a virus as given above in combination with a pharmaceutically acceptable carrier. The formulation is useful for both raising antibodies in animals, which antibodies specifically bind to the virus and are useful in diagnostic assays and other methods of detecting the virus in both humans and animals; the formulation is useful as a vaccine formulation for producing protective immunity against the virus in an animal or in a human subject.

A fourth aspect of the present invention is a method of producing an immune response (e.g., producing antibodies and/or producing protective immunity) in a subject. The method comprises administering a virus as given above to the subject in an amount effective to produce an immune response in that subject.

A fifth aspect of the present invention is the use of a virus as described above for the preparation of a medicament for producing an immune response in a subject, as described above.

A sixth aspect of the present invention is a method of making an attenuated virus as given above. The method comprises providing a host cell containing an expression vector, the expression vector containing a DNA encoding the attenuated virus, which host cell does not methylate the DNA sufficiently to block the expression of the viral DNA; and expressing the attenuated virus in said host cell. Typically, the host cell is provided in a suitable incubation media, the virus collected from the media after expression therein (with lysis of the host cell, if necessary), and the media either used directly

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to produce an immune response in a subject, or the virus collected and/or purified from the media and then combined with other ingredients to produce a pharmaceutical formulation.

The foregoing and other objects and aspects of the present invention are explained in detail in the specification set forth below.

Brief Description of the Drawings

Figure 1 illustrates the interruption of the 10 life-cycle of CpG-inserted retrovirus genomes.

Figure 2 illustrates the CpG content of HIV-1 strain HIVHX2CG (F. Wong-Staal et al., Nature 313, 277-284 (1985).

Figure 3 illustrates the CpG content of an HIV-15 1 genome of the present invention, strain HIV-1^{CpG1} (SEQ ID NO:1).

Detailed Description of the Invention

The nucleotide sequence of an HIV-1 genome (HIV-1^{cpG1}) modified according to the principles described 20 herein is presented by single strand only, in the 5' to 3' direction, from left to right. Nucleotides and amino acids are represented herein in the manner recommended by the IUPAC-IUB Biochemical Nomenclature Commission, or (for amino acids) by three letter code, in accordance with 37 CFR §1.822 and established usage. See, e.g., PatentIn User Manual, 99-102 (Nov. 1990) (U.S. Patent and Trademark Office). In nucleotide sequences herein, the internucleotide phosphate linkage is sometimes designated with a "p" positioned between the standard single capital letter for the nucleotide, as in "CpG" for 5'-CG-3'.

1. Viruses

The viruses of the present invention are, in general, expression defective viruses. That is, for the purpose of manufacturing the virus, the virus genome or a DNA encoding the virus genome may be introduced into a host cell that does not methylate the viral DNA sufficient to inactivate it. The viral genome can thus be

transcribed into RNA in such a host cell, the RNA then translated into viral proteins, and encapsidated viral genomes (viral particles) produced. For the purpose of producing an immune response in an animal or human 5 subject, the target cells in this case do methylate the viral genome such that methylation sensitive processing of the viral genome, such as transcription, is inhibited The present invention may accordingly be therein. carried out with any virus in which the genome of the 10 virus is methylated in the cells of the subject to which the virus is administered, including DNA viruses, RNA viruses and retroviruses. Retroviruses are particularly preferred. A schematic of the life cycle of a retrovirus and an illustration of how CpG-inserted retrovirus genomes interrupt the life cycle is given in Figure 1. Note that in Figure 1, stages of the life cycle depicted interrupted in CpG inserted by bold lines are retrovirions.

Retroviruses that may be used to carry out the 20 present invention include retroviruses of both animals and man. This group of retroviruses includes both simple retroviruses and complex retroviruses. The of B-type the subgroups retroviruses include retroviruses and D-type retroviruses. C-type 25 retroviruses. An example of a B-type retrovirus is mouse mammary tumor virus (MMTV). The C-type retroviruses include subgroups C-type group A (including Rous sarcoma virus (RSV), avian leukemia virus (ALV), and avian myeloblastosis virus (AMV)) and C-type group B (including 30 murine leukemia virus (MLV), feline leukemia virus (FeLV), murine sarcoma virus (MSV), gibbon ape leukemia necrosis virus (GALV). spleen reticuloendotheliosis virus (RV) and simian sarcoma virus The D-type retroviruses include Mason-Pfizer 35 monkey virus (MPMV) and simian retrovirus type 1 (SRV-1). The complex retroviruses include the subgroups lentiviruses, T-cell leukemia viruses and the foamy

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viruses. Lentiviruses include HIV-1, but also include HIV-2, SIV, Visna virus, feline immunodeficiency virus (FIV), and equine infectious anemia virus (EIAV). The T-cell leukemia viruses include HTLV-1, HTLV-II, simian T-cell leukemia virus (STLV), and bovine leukemia virus (BLV). The foamy viruses include human foamy virus (HFV), simian foamy virus (SFV) and bovine foamy virus (BFV). The foregoing is illustrative, and is not intended to be limiting of the retroviruses that may be employed in carrying out the instant invention.

Examples of other RNA viruses that may be used in carrying out the present invention include, but are not limited to, the following: members of the family Reoviridae, including the genus Orthoreovirus (multiple serotypes of both mammalian and avian retroviruses), the qenus Orbivirus (Bluetongue virus, Eugenangee virus, Kemerovo virus, African horse sickness virus, Colorado Tick Fever virus), the genus Rotavirus (human calf diarrhea rotavirus. Nebraska virus, rotavirus, simian rotavirus, bovine or ovine rotavirus, 20 avian rotavirus); the family Picornaviridae, including the genus Enterovirus (poliovirus, Coxsackie virus A and B, enteric cytopathic human orphan (ECHO) viruses, virus, Simian enteroviruses, hepatitis Α encephalomyelitis (ME) viruses, Poliovirus muris, Bovine enteroviruses, Porcine enteroviruses), the genus Cardiovirus (Encephalomyocarditis virus (EMC), Mengovirus), the genus Rhinovirus (Human rhinoviruses at least 113 subtypes; other rhinoviruses), the genus Apthovirus (Foot and Mouth disease (FMDV); the family 30 Calciviridae, including Vesicular exanthema of swine virus, San Miquel sea lion virus, Feline picornavirus and Norwalk virus; the family Togaviridae, including the genus Alphavirus (Eastern equine encephalitis virus, Semliki forest virus, Sindbis virus, Chikungunya virus, 35 O'Nyong-Nyong virus, Ross river virus, Venezuelan equine encephalitis virus, Western equine encephalitis virus),

the genus Flavirius (Mosquito borne yellow fever virus, Dengue virus, Japanese encephalitis virus, St. Louis encephalitis virus, Murray Valley encephalitis virus, West Nile virus, Kunjin virus, Central European tick 5 borne virus, Far Eastern tick borne virus, Kyasanur forest virus, Louping III virus, Powassan virus, Omsk hemorrhagic fever virus), the genus Rubivirus (Rubella virus), the genus Pestivirus (Mucosal disease virus, Hog the cholera virus, Border disease virus); 10 Bunyaviridae, including the genus Bunyvirus (Bunyamwera California encephalitis and related viruses, viruses), the genus Phlebovirus (Sandfly fever Sicilian virus, Rift Valley fever virus), the genus Nairovirus (Crimean-Congo hemorrhagic fever virus, Nairobi sheep 15 disease virus), and the genus Uukuvirus (Uukuniemi and related viruses); the family Orthomyxoviridae, including the genus Influenza virus (Influenza virus type A, many human subtypes); Swine influenza virus, and Avian and Equine Influenza viruses; influenza type B (many human and influenza type C (possible separate 20 subtypes), genus); the family paramyxoviridae, including the genus Paramyxovirus (Parainfluenza virus type 1, Sendai virus, Hemadsorption virus, Parainfluenza viruses types 2 to 5, Newcastle Disease Virus, Mumps virus). the subacute sclerosing 25 Morbillivirus (Measles virus, distemper virus, Rinderpest panencephalitis virus, virus), the genus Pneumovirus (respiratory syncytial virus (RSV), Bovine respiratory syncytial virus and Pneumonia virus of mice); the family Rhabdoviridae, 30 including the genus Vesiculovirus (VSV), Chandipura virus, Flanders-Hart Park virus), the genus Lyssavirus (Rabies virus), two genera of fish Rhabdoviruses, and two probable Rhabdoviruses (Marburg virus and Ebola virus); Arenaviridae, including Lymphocytic family the 35 choriomeningitis virus (LCM), Tacaribe virus complex, and family Coronoaviridae, the Lassa virus; Infectious Bronchitis Virus (IBV), Mouse Hepatitis virus,

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Human enteric corona virus, and Feline infectious peritonitis (Feline coronavirus).

Illustrative DNA viruses that may be employed in carrying out the present invention include, but are 5 not limited to: the family Poxviridae, including the genus Orthopoxvirus (Variola major, Variola minor, Monkey pox Vaccinia, Cowpox, Buffalopox, Rabbitpox, Ectromelia), the genus Leporipoxvirus (Myxoma, Fibroma), the genus Avipoxvirus (Fowlpox, other avian poxvirus), the genus 10 Capripoxvirus (sheeppox, goatpox), the genus Suipoxvirus (Swinepox), the genus Parapoxvirus (contagious postular dermatitis virus, pseudocowpox, bovine papular stomatitis virus); the family Iridoviridae (African swine fever virus, Frog viruses 2 and 3, Lymphocystis virus of fish; Herpesviridae, including the family 15 the Herpesviruses (Herpes Simplex Types 1 and 2, Varicella-Zoster, Equine abortion virus, Equine herpes virus 2 and pseudorabies virus, infectious bovine bovine virus, infectious keratoconjunctivitis feline rhinotracheitis virus, 20 rhinotracheitis virus, laryngotracheitis Betavirus; the infections (Human cytomegalovirus and herpesviruses cytomegaloviruses of swine, monkeys and rodents; the gamma-herpesviruses (Epstein-Barr virus (EBV), Marek's 25 disease virus, Herpes saimiri, Herpesvirus ateles, Herpesvirus sylvilagus, guinea pig herpes virus, Lucke tumor virus; the family Adenoviridae, including the genus Mastadenovirus (Human subgroups A, B, C, D, E and ungrouped; simian adenoviruses (at least 23 serotypes), infectious 30 canine hepatitis, and adenoviruses of cattle, pigs, frogs and many other species, the sheep, Aviadenovirus (Avian adenoviruses); and non-cultivatable adenoviruses; the family Papoviridae, including the genus Papillomavirus (Human papilloma viruses, many subtypes, 35 bovine papilloma viruses, Shope rabbit papilloma virus, and various pathogenic papilloma viruses of other species), the genus Polyomavirus (polyomavirus, Simian

vacuolating agent (SV-40), Rabbit vacuolating agent (RKV), K virus, BK virus, JC virus, and other primate polyoma viruses such as Lymphotrophic papilloma virus; the family Parvoviridae including the genus Adeno-Parvovirus (Feline 5 associated viruses, the qenus canine virus, bovine parvovirus, panleukopenia parvovirus, Aleutian mink disease virus, etc). Finally, DNA viruses may include viruses which do not fit into the above families: Kuru, Creutzfeldt-Jacob disease viruses 10 and chronic infectious neuropathic agents (CHINA virus).

2. Introduction of Attenuating Mutations

The phrase "attenuated virus", as used herein, means that the infection of a susceptible host by that virus will result in decreased probability of causing 15 disease in its host (loss of virulence) in accord with standard terminology in the art. See, e.g, B. Davis, R. Dulbecco, H. Eisen, and H. Ginsberg, Microbiology, 132 (3rd ed. 1980). Attenuating mutations are mutations that cause a virus that would otherwise be capable of causing Viruses of the 20 disease to be an attenuated virus. instant invention are attenuated in the sense that the viral life cycle in the susceptible host is inhibited at the level of transcription for retroviruses and DNA viruses. In the case of non-retroviral RNA viruses, the 25 viral life cycle is assumed to be inhibited by loss of as a result of CpG the RNA genome function of methylation.

The number of additional methylation sites introduced by mutation of the genome of a virus as given above to produce a modified virus of the invention may be relatively few (e.g., 1, 2, or 3), or may be at least 10, 50, 100 or 500 or more, depending on the site of the mutation, the nature of the virus, the presence or absence of other attenuating mutations (e.g., a deletion of the nef gene in a retrovirus), etc. Typically, a sufficient number of methylation sites are introduced into the genome of the virus so that the ratio of

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observed to expected CpG dinucleotides (CpG^{o/e}) within the genome will be increased over that found in the wild type virus 1, 2, 3, 4, 5, 6, 7 or 8-times or more, though the increase in CpG^{o/e} need not be increased as much where a few methylation sites that are particularly active as attenuating mutations are employed.

Modified viruses of the present invention are, in general, infectious virus particles comprising a viral capsid containing the nucleic acid material (DNA or RNA)

10 that comprises the viral genome, which particles bind to the target cells in the subject to which they are administered and introduce their genome into those cells. It is accordingly preferred that the modified virus contain at least two or three mutations that are attenuating (whether by the introduction of a methylation site as described herein or by another mechanism) to reduce the possibility of the virus spontaneously reverting to virulence.

Attenuating CpG mutations of the instant invention are introduced into cDNAs encoding virus by any suitable means, such as by direct synthesis, PCR mutagenesis, or site-directed mutagenesis (see, e.g., U.S. Patent No. 4,873,192 to Kunkel) (applicant specifically intends that the disclosure of all patent references cited herein be incorporated herein by reference).

The attenuated viruses of the present invention are produced directly on a DNA synthesizing machine, the use of which is known in the art. Specifically, the nucleic acid sequence of the target virus (for example, HIV-1) is selected. The genome is then scanned for non-CpG containing codons which have the possibility of being changed to CpG-containing codons without altering the resulting post-translational amino acid sequence. These non-CpG-containing codons are thus replaced with CpG dinucleotides. For example, a proline coded for by CCT, CCC, or CCA would be switched to CCG. Alternatively,

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adjacent codons are altered such that they contain a CpG within their adjoining region. As an example, the adjacent codons GCA GTG (alanine-valine) can be altered to GCC GTG, which still codes for alanine-valine but now contains a methylatable CpG (the last C of the first codon and the beginning G of the second).

Of course, certain codons are preferred over others in a species-specific way. It is preferable to create altered genomes by selecting preferred codons where possible (i.e., codons preferred in both the host cell culture system in which the virus is produced, and codons preferred in the subject administered the virus to produce an immune response therein).

Viruses of the present invention can, as noted include additional attenuation strategies in 15 addition to the inclusion of the silent CpG mutations example, a conventional For described herein. substitution mutation that produces an amino acid substitution that is attenuating in the encoded protein 20 may also be included, if desired. As another example using HIV-1, the nef gene and another gene or genes or portions thereof can be deleted so as to produce attenuating mutations thereof.

In the case of a retrovirus such as HIV-1, in which many strains of the virus are present, it may be desirable to modify multiple HIV-1 strains by CpG insertion, using them together to produce an effective vaccine.

A novel HIV-1 genome (hereinafter referred to has been hypersubstituted with HIV-1^{CpG-1}) that 30 as CpGs "silent" noninformational or Non-informational means that addition of hereinbelow. the CpGs to the genome does not alter the amino acid The proteins. resulting in the sequence 35 hypersubstitution of CpGs makes this novel synthetic genome a target for host cell methylases. Thus, although the virus for which this genome codes is capable of

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infecting the cell, the proviral genome is easily inactivated by methylation and kept permanently in a dormant state. That is, to the extent the genome can be methylated by the host, it will remain transcriptionally 5 silent.

While the present invention is contemplated primarily for use with so-called "live" virus vaccines, it may also be used with killed virus vaccines, including formaldehyde and heat-inactivated viruses. 10 invention is useful in such vaccine preparations because occasionally live virus escapes the killing procedure and can cause infection. Thus the instant invention, used in conjunction with any other attenuation strategy, provides a further level of attenuation.

15 3. Production of Virus in Cell Culture

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An expression vector is a replicable DNA construct in which a DNA sequence encoding one or more proteins is operably linked to suitable control sequences capable of affecting the expression of the DNA in a 20 suitable host. A replication vector may be used to produce additional DNA where expression of that DNA is Choice of host cell for a particular not necessary. vector will depend upon factors such as whether expression or replication is desired.

Transformed host cells are cells which have been transformed or transfected with vectors constructed using recombinant DNA techniques. Transformed host cells ordinarily express the DNA, but host cells transformed for purposes of cloning or amplifying the target proteins 30 do not need to express the protein.

cells Suitable host generally prokaryote, yeast or higher eukaryotic cells such as mammalian cells and insect cells. Cells derived from multicellular organisms are a particularly suitable host for recombinant methylated viruses, and insect cells are Propagation of such cells in particularly preferred. cell culture has become a routine procedure (Tissue -13-

Culture, Academic Press, Kruse and Patterson, editors (1973)). Examples of useful host cell lines are CD4+ T lymphocytes such as MOLT4, VERO and HeLa cells, Chinese hamster ovary (CHO) cell lines, and WI138, BHK, COS-7, 5 CV, and MDCK cell lines. Expression vectors for such cells ordinarily include (if necessary) an origin of replication, a promoter located upstream from the DNA encoding the methylatable virus to be expressed and operatively associated therewith, along with a ribosome binding site, an RNA splice site (if intron-containing genomic DNA is used), a polyadenylation site, and a transcriptional termination sequence.

Where the host cell contains a methylation system that would otherwise methylate the viral genome, 15 that methylation system must be inactivated sufficiently to permit production of the virus therein. Such inactivation may be accomplished by any suitable means, such as by including a demethylating agent or methylase inhibitor such as 5-azacytidine or 5-azadeoxycytidine in 20 the cell culture media in an amount sufficient to inhibit the methylation system (e.g., 1-10 μ M), by adding an antisense oligonucleotide to the media in an amount effective to inactivate the methylation system, or by genetically engineering the cells to express an antisense 25 agent therein effective to inactivate the methylation the antisense system is genetically Where system. engineered into the cell, it is most preferable to use an inducible expression vector, for example on in which the antisense oligonucleotide is placed downstream of a 30 promoter such as the mouse metallothionein promoter, which can be activated to express the antisense by addition of a metal (such as cadmium) to the tissue Numerous such inducible expression culture medium. systems are known to those skilled in the art.

25 Expressing live virus is particularly feasible in a Baculovirus expression system, which utilizes insect cells as the host cells and viral vectors indigenous to

insects (See generally U.S. Patents Nos. 4,745,051 and 4,879,236 to Smith et al.). Baculoviruses are members of the family Baculoviridae and the genus Baculovirus. genus comprises three subgroups of viruses: the nuclear 5 polyhedrosis viruses (NPV), the granulosis viruses (GV) and the non-occluded viruses. NPVs include Autographica californica NPV (AcNPV), Heliothis zea NPV (HzNPV) and Bombyx mori NPV (BmNPV). The use of recombinant baculovirus vectors to express foreign proteins in insect 10 cell cultures or larvae is known. See e.g., Luckow & Summers, Bio/Technology, 6, 47 (1988); Tomalski & Miller, Nature, 352, 82 (1991). The use of baculoviruses in this invention is particularly useful because insect host cells (e.g., cultured Spodoptera frugiperda cells) do not 15 possess DNA methylase enzymes and cannot therefore transcriptionally inactivate the viral proviral DNA. general, a baculovirus expression vector comprises a baculovirus genome containing the DNA to be expressed inserted into the polyhedrin gene at a position where it 20 is under the transcriptional control of the baculovirus polyhedrin promoter.

Modified virus produced by tissue culture techniques as described above can be isolated and/or purified as desired by techniques such as ultrafiltration, and then combined with other ingredients to provide the modified virus in a pharmaceutically acceptable carrier.

4. Pharmaceutical Formulations

composition of matter comprising 30 preparation of the attenuated viral particles produced by the cell line of the present invention is disclosed include This composition may any herein. pharmaceutically acceptable carrier (such as sterile, pyrogen-free physiological saline solution, or sterile, 35 pyrogen-free phosphate-buffered saline solution). general, the compositions are prepared by contacting and combining viral particles produced as above with a

pharmaceutically acceptable carrier. The viral particles of the composition may be live, killed, fixed or lyophilized, as is most suitable for the intended use. The viral particles are included in the composition in an immunogenic amount, the amount to be determined by the intended use. The immunogenic activity of a given amount of the virus of the present invention may be determined by any of a number of methods known in the art. The increase in titer of antibody against a particular viral antigen upon administration may be used as a criteria for immunogenic activity.

Subjects which may be administered the live attenuated viruses and formulations disclosed herein include both human subjects and animal subjects (e.g., the veterinary treatment of primates such as owl monkeys, marmosets and chimpanzees, and other mammalian species such as dogs, cats, pigs, and horses, and non-mammalian species such as birds (chickens, turkeys, etc.)).

Pharmaceutical formulations of the present invention comprise an immunogenic amount of a live attenuated virus as disclosed herein in combination with a pharmaceutically acceptable carrier. An "immunogenic amount" is an amount of the attenuated virus sufficient to evoke an immune response in the subject to which the virus is administered. The particular dose employed is not critical, and depends upon the type and condition of the subject, the route of administration, etc.

Techniques to determine a particular immunogenic amount of the viral particles of the present invention will be apparent to those of ordinary skill in the art. For example, the active agent (viral particles or preparations thereof) may be given in an amount of from .01 to 100 μ g per Kg body weight (e.g., .5 or 1.0 μ g per Kg).

Administration of the live attenuated viruses disclosed herein may be carried out by any suitable means, including both parenteral injection (such as

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intramuscular intraperitoneal, subcutaneous, or injection), by oral administration, and by topical application of the virus (typically carried in the pharmaceutical formulation) to an airway surface. 5 Topical application of the virus to an airway surface can be carried out by intranasal administration (e.g., by use of a dropper, swab, or inhaler which deposits a formulation intranasally). pharmaceutical application of the virus to an airway surface can also be 10 carried out by inhalation administration, such as by creating respirable particles of a pharmaceutical formulation (including both solid particles and liquid particles) containing the virus as an aerosol suspension, and then causing the subject to inhale the respirable 15 particles. Methods and apparatus for administering respirable particles of pharmaceutical formulations are and any conventional technique can be employed. See, e.g., U.S. Patent No. 5,304,125 to D. Leith; U.S. Patent No. 5,299,566 to C. Davis and R. 20 Snyder; U.S. Patent No. 5,290,550 to R. Fisher and W. Metzger; and U.S. Patent No. 5,292,498 to R. Boucher.

Oral vaccine formulations may be made from a culture of cells producing live virus containing the desired attenuating mutations in accordance with known techniques. The culture itself may be administered to the subject; the culture may be optionally filtered and/or clarified; stabilizers such as sucrose, MgCl₂, etc. may be added to the media. Pharmaceutically acceptable carriers for oral administration may be a syrup, elixir, lozenge, etc. The vaccine formulation may be prepared in accordance with known techniques, such as illustrated by R. Purcell et al., Vaccine Against Hepatitis A Virus, U.S. Patent No. 4,894,228.

While the viruses, methods and formulations of 35 the present invention have been described above with reference to producing protective immunity by the administration of vaccine formulations, they may also be

used to immunize animals to simply produce antibodies in animals, which antibodies may then be collected and used for the purpose of detecting and/or diagnosing various viral infections or the presence of viral particles in 5 biological samples in accordance with conventional See generally E. Maggio, Enzyme diagnostic techniques. Immunoassay (1980); see also U.S. Patents Nos. 4,659,678, 4.376,110, 4,275,149, 4,233,402, and 4,230,767.

5. Oligonucleotide probes

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An advantage of the instant invention is that it will permit detection of infection by wild-type virus even after vaccination has occurred. For example, a vaccine employing a whole or nearly whole virus will create an immune response to the virus that will preclude 15 standard immunologic or nucleic acid detection The constructs of the instant subsequent infection. invention, since they represent totally new creations at the level of the DNA, can easily be distinguished by molecular probing. Thus, probes can be made that will be 20 specific for the wild type virus and that will not hybridize to a virus of the instant invention, and probes can be made that will specifically bind to the virus of the instant invention and not the wild type virus.

Thus, a further aspect of the present invention 25 is an oligonucleotide probe useful for distinguishing between (i) an attenuated virus, not naturally occurring, containing at least 1 additional methylation site in the genome of the virus compared to the corresponding wildtype virus, and (ii) the corresponding wild-type virus, 30 with the oligonucleotide probe selected from the group oligonucleotide probes consisting of: (a) selectively hybridize to the nucleic acid of attenuated virus of (i) above, and which do not hybridize to the nucleic acid of the wild-type virus of (ii) above 35 under the same hybridization conditions; oligonucleotide probes that selectively hybridize to the nucleic acid of a wild-type virus of (ii) above,

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which do not hybridize to the nucleic acid of the of (i) above under the same attenuated virus The probe may be of any hybridization conditions. suitable length so long as the desired specificity of 5 binding is achieved. Such probes are typically at least 8 to 12 nucleotides in length and can be up to 20-40 nucleotides or more in length. The probe may be of any suitable nucleic acid, including DNA and RNA. The probe may be labeled with or conjugated to a detectable group 10 (e.g., a radioisotope such as ³²P, ¹²⁵I, ¹³¹I, ³H, ¹⁴C, or 35S; an enzyme such as horseradish peroxidase or alkaline phosphatase) by a variety of techniques, including direct The probe may be one probe or a member covalent bond. a pair of probes useful for a nucleic acid 15 amplification procedure, such as polymerase chain reaction (PCR), ligase chain reaction (LCR), or strand displacement amplification (SDA). Techniques for use of such probes are known to those skilled in the art. See, for example, U.S. Patent No. 4,358,535 to Falkow and 20 Mosley; U.S. Patent No. 4,302,204 to Wahl and Stark; U.S. Patent No. 4,994,373 to Stavrianopoulos; U.S. Patent No. 5,270,184 to Walker et al.; and, for PCR, U.S. Patents Nos. 4,683,195, 4,683,202, 4,800,159 and 4,965,188.

The present invention is explained in greater 25 detail in the following non-limiting Examples.

EXAMPLE 1

Introduction of CpG Sites in HIV-1 Genome

The genomic sequence of HIV-1 strain HIVHXB2CG (see, e.g., F. Wong-Staal et al., Nature 313, 277-284 (1985) was obtained from GENBANK (Accession Number k03445). Sites in the sequence in which silent substitution mutations could be added to the genome to introduce additional CpG segments therein were identified and a new DNA encoding a non-natural derivative of the HIV-1 genome is synthesized as follows.

Single stranded DNA segments 75 bases in length are synthesized by phosphoramidate chemistry on an

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Applied Biosystems Model 394 DNA/RNA Synthesizer (Applied Biosystems Inc., 850 Lincoln Centre Drive. Foster City, California, 94404 USA). Each 75 base pair double-stranded DNA segment is deprotected at 55°C for 12 hours and dried to remove ammonium hydroxide. The trityl group is left on at the deprotecting step. The full-length 75 base-pair segment is then separated from shorter "failure" segments in the preparation with NENSORB™ chromatography. This serves to avoid adding the shorter failure segments to the elongated segment.

Complementary segments are made and annealed together, with overlapping ends of 4 bases, to produce a double-stranded DNA segment 75 bases in length. Each new 75 base-pair double-stranded segment is sequentially ligated to the previous segment to build up an elongated double-stranded DNA segment that ultimately becomes the entire modified HIV-1 genome (HIV-1^{cpG-1}), given in SEQ ID NO:1.

Appropriate splice segments are added to each 20 end of the complete genome by conventional techniques and the genome inserted into an expression vector.

COMPARATIVE EXAMPLE A

Comparison of CpG Sites in HIV-1 Strain HIVHXB2CG and Strain HIV-1^{cpG1}

The CpG content of the HIVHXB2CG genome is illustrated in graph form in Figure 2. The gene structure of HIV is incorporated into this graph for clarity.

The CpG content of the HIV-1^{CpG1} genome is illustrated by graph in Figure 3. Note the dramatic increase in CpG content as compared to the wild-type genome shown in Figure 2. HIV-1^{CpG1} has 948 new CpG sites as compared to HIVHXB2CG (representing a more than tenfold increase in CpG segments: 97 in HIVHXB2CG; 1045 in HIV-1^{CpG1}). The ratio of expected over observed CpG dinucleotides (CpG^{O/e} in HIV-1^{CpG-1} is increased from a value of 0.22 in HIVHXB2CG to a value of 1.68 in HIV-1^{CpG-1}.

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This represents an approximately 8-fold increase in CpG^{o/e}. In extreme cases (e.g. those in which many hundreds of new CpG methylation sites have been inserted into the viral genome, as in the example modified genome, HIV-1^{cpG-1}) this will result in an increase in the GC/CT ratio above that observed in the wild type virus. Thus, the GC/AT ratio in HIV-1^{cpG-1} is equal to 1.05 as compared to 0.74 in the wild type genome, HIVHXB2CG. The base count in HIV-1^{cpG-1} as compared to HIVHXB2CG is as follows:

		HIVHXB2CG	HIV-1 ^{CpG-1}
10	Adenines	3411	2796
:	Cytosines	1773	2197
	Guanines	2370	2772
	Thymines	2164	1953

This represents a loss of 615 adenines and 211 thymines in HIV-1^{CpG-1} as compared to HIVHXB2CG and a gain of 424 cytosines and 402 guanines in HIV-1^{CpG-1} as compared to HIVHXB2CG. The ration of GC/AT will not be increased significantly in those modified genomes in which only a small number of CpGs need to be inserted (e.g. < 10) to interrupt the viral life cycle. The GC/AT ratio in HIVHXB2CG is 0.74; while the GC/AT ratio in HIV-1^{CpG1} is 1.05.

EXAMPLE 2

Expression of HIV-1 Genome in Insect Cells

The BACKPACK™ baculovirus expression system is obtained from Clontech Inc. (Telephone Number in USA: 415-424-8222). The genomic DNA segment described in Example 1 above is ligated into the multiple cloning site of pBacPAK8™ (or PBacPAK9™) to produce a recombinant vector, with expression of the genomic DNA driven by the strong AcMNPV polyhedrin promoter in the vector.

10 Cultured Spodoptera frugiperda cells are transformed with the recombinant vector and the virus of the invention is produced in the cultured cells in accordance with the manufacturer's instructions.

The foregoing examples are illustrative of the present invention, and are not to be construed as limiting thereof. The invention is defined by the following claims, with equivalents of the claims to be included therein.

SEQUENCE LISTING

- (1) GENERAL INFORMATION:
 - (i) APPLICANT: Nyce, Jonathan W.
 - (ii) TITLE OF INVENTION: Attenuated Viruses and Method of Making the Same
 - (jii) NUMBER OF SEQUENCES: 1
 - (iv) CORRESPONDENCE ADDRESS:
 (A) ADDRESSEE: Kenneth D. Sibley (B) STREET: Post Office Box 34009 (C) CITY: Charlotte (D) STATE: North Carolina

 - (E) COUNTRY: USA (F) ZIP: 28234

 - (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk

 - (B) COMPUTER: IBM PC compatible
 (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
 - (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 08/319.974
 - (B) FILING DATE: 07-OCT-1994
 - (C) CLASSIFICATION:
 - (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Sibley, Kenneth D.
 - (B) REGISTRATION NUMBER: 31,665
 - (C) REFERENCE/DOCKET NUMBER: 5218-27
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: 919-881-3140 (B) TELEFAX: 919-881-3175 (C) TELEX: 575102
- (2) INFORMATION FOR SEQ ID NO:1:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9718 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

60	ATCTACCACA	TGATCTGTGG	AAGATATCCT	CAACGAAGAC	AATTCACTCC	TGGAAGGGCT
120	AGATATCCAC	GCCAGGGATC	ACACACCAGG	TAGCAGAACT	CTTCCCTGAT	CACAAGGCTA
180	GAAGAAGCCA	AGAGAAGTTA	CAGTTGAGCC	AAGCTAGTAC	ATGGTGCTAC	TGACCTTTGG
240	GATGACCCGG	GCATGGAATG	CTGTGAGCCT	TTGTTACACC	GAACACCAGC	ACAAAGGAGA
300	ATGGCCCGAG	ATTTCATCAC	GCCGCCTAGC	AGGTTTGACA	GTTAGAGTGG	AGAGAGAAGT
360	GGACTTTCCG	TTGCTACAAG	GACATCGAGC	AAGAACTGCT	GGAGTACTTC	AGCTGCATCC
420	GCCCTCAGAT	GGAGTGGCGA	GCGGGACTGG	CGTGGCCTGG	TCCAGGGAGG	CTGGGGACTT
480	CCAGATCTGA	TCTGGTTAGA	ACTGGGTCTC	TTTTGCCTGT	AGCAGCTGCT	CCTGCATATA
540	AAGCTTGCCT	AGCCTCAATA	CCACTGCTTA	ACTAGGGAAC	TCTCTGGCTA	GCCTGGGAGC
600	GAGATCCCTC	CTGGTAACTA	TTGTGTGACT	TGCCCGTCTG	AAGTAGTGTG	TGAGTGCTTC
660	GACCTGAAAG	CCCGAACAGG	AGCAGTGGCG	GAAAATCTCT	AGTCAGTGTG	AGACCCTTTT
720	CCCGCACGGC	TGCTGAAGCG	GGACTCGGCT	TCTCGACGCA	ACCAGAGCTC	CGAAAGGGAA
780	GAGGCTAGAA	TTGACTAGCG	GCCAAAAATT	TGGTGAGTAC	GGGCGGCGAC	AAGAGGCGAG
840	CGATGGGAAA	CGAATTAGAT	TATCGGGCGG	GCGTCGGTAT	GGGCGCGCGC	GGAGAGAGAT
900	GTATGGGCGT	AAAACATATC	AATATAAATT	GGAAAGAAAA	ACGGCCGGGC	AAATTCGGTT
960	GAAGGCTGTC	AGAAACGTCG	CGGGCCTGTT	GCGGTTAATC	CGAACGATTC	CGCGCGAGCT
1020	CTTCGATCGT	ATCGGAAGAA	TTCAGACGGG	CAACCGTCGC	CGGACAGCTA	GACAAATACT
1080	AAAGACACGA	GATCGAGATA	TCCATCAACG	CTCTATTGCG	GGTCGCGACG	TATATAATAC
1140	GCGCAGCAAG	GAAGAAAAA	AAAACAAATC	GAGGAAGAGC	AGACAAGATC	AGGAAGCGTT
1200	GTGCAGAACA	TTACCCGATC	TCTCGCAAAA	TCGAATCAGG	CACGGGACAC	CGGCGGCGGA
1260	TGGGTAAAAG	GTTAAACGCG	CGCCGCGAAC	CAGGCGATAT	AATGGTACAT	TCCAGGGGCA
1320	TTATCGGAAG	GTTTTCGGCG	TGATACCGAT	TCGCCGGAAG	GAAGGCGTTC	TCGTCGAAGA
1380	CAAGCGGCGA	CGGCGGACAT	TAAACACGGT	AACACGATGC	GCAAGATTTA	GAGCGACGCC
1440	GTGCATCCGG	ATGGGATCGC	AAGCGGCGGA	ATCAACGAGG	AAAAGAGACG	TGCAAATGTT
1500	GACATCGCGG	GCGCGGATCG	TGCGCGAACC	CCGGGCCAGA	GCCGATCGCG	TGCACGCGG
1560	CCGATCCCGG	GAATAATCCG	GATGGATGAC	GAACAAATCG	GACGCTTCAG	GAACGACGTO
1620	CGAATGTATT	TAAAATCGTA	TCGGATTAAA	TGGATAATCC	TTATAAACGA	TCGGCGAAA
1680	GACTACGTCG	ACCGTTTCGC	GACCGAAGGA	ATACGACAAG	GATTCTCGAC	CGCCGACGT

ACCGGTTCTA	TAAAACGCTA	CGCGCGGAGC	AAGCGTCGCA	GGAGGTAAAA	AATTGGATGA	1740
CGGAAACGTT	GTTGGTCCAA	AACGCGAACC	CGGATTGTAA	GACGATTTTA	AAAGCGTTGG	1800
GACCGGCGGC	GACGCTCGAA	GAAATGATGA	CGGCGTGTCA	GGGCGTCGGC	GGACCGGGCC	1860
ATAAGGCGCG	CGTTTTGGCG	GAAGCGATGT	CGCAAGTAAC	GAATTCGGCG	ACGATAATGA	1920
TGCAGCGCGG	CAATTTTCGG	AACCAACGAA	AGATCGTTAA	GTGTTTCAAT	TGCGGCAAAG	1980
AAGGGCACAC	GGCGCGAAAT	TGCCGCGCGC	CGCGGAAAAA	GGGCTGTTGG	AAATGCGGAA	2040
AGGAAGGACA	CCAAATGAAA	GATTGTACGG	AGCGACAGGC	GAATTTTCTC	GGGAAGATCT	2100
GGCCGTCGTA	CAAGGGACGG	CCGGGGAATT	TTCTTCAGTC	GCGACCGGAG	CCGACGGCGC	2160
CGCCGGAAGA	GTCGTTCCGG	TCGGGCGTCG	AGACGACGAC	GCCGCCGCAG	AAGCAGGAGC	2220
CGATAGACAA	GGAACTGTAT	CCGTTAACGT	CGCTCCGGTC	GCTCTTCGGC	AACGACCCGT	2280
CGTCGCAATA	AAGATAGGGG	GGCAACTAAA	GGAAGCTCTA	TTAGATACAG	GAGCAGATGA	2340
TACAGTATTA	GAAGAAATGT	CGTTGCCGGG	ACGATGGAAA	CCGAAAATGA	TCGGCGGAAT	2400
CGGCGGTTTT	ATCAAAGTAC	GACAGTACGA	TCAGATACTC	ATCGAAATCT	GCGGACATAA	2460
AGCGATCGGT	ACCGTACTCG	TCGGACCGAC	GCCGGTCAAC	ATAATCGGAC	GAAATCTGTT	2520
GACGCAGATC	GGTTGCACGT	TAAATTTTCC	GATTTCGCCG	ATCGAGACGG	TACCGGTAAA	2580
ATTAAAGCCG	GGAATGGACG	GCCCGAAAGT	TAAACAATGG	CCGTTGACCG	AAGAAAAAAT	2640
AAAAGCGCTC	GTCGAAATTT	GTACGGAGAT	GGAAAAGGAA	GGGAAAATTT	CGAAAATCGG	2700
GCCGGAAAAT	CCGTACAATA	CGCCGGTATT	CGCGATAAAG	AAAAAAGACT	CGACGAAATG	2760
GCGAAAACTC	GTCGATTTCC	GCGAACTTAA	TAAGCGAACG	CAAGACTTCT	GGGAAGTTCA	2820
ACTCGGAATA	CCGCATCCGG	CGGGGTTAAA	AAAGAAAAA	TCGGTAACGG	TACTCGATGT	2880
CGGCGACGCG	TATTTTTCGG	TTCCGCTCGA	CGAAGACTTC	CGGAAGTATA	CGGCGTTTAC	2940
GATACCGTCG	ATAAACAACG	AGACGCCGGG	GATTCGATAT	CAGTACAACG	TGCTTCCGCA	3000
GGGATGGAAA	GGATCGCCGG	CGATATTCCA	ATCGTCGATG	ACGAAAATCC	TCGAGCCGTT	3060
TCGAAAACAA	AATCCGGACA	TCGTTATCTA	TCAATACATG	GACGATTTGT	ACGTCGGATC	3120
GGACCTCGAA	ATCGGGCAGC	ATCGAACGAA	AATCGAGGAG	CTGCGACAAC	ATCTGTTGCG	3180
GTGGGGACTT	ACGACGCCGG	ACAAAAAACA	TCAGAAAGAA	CCGCCGTTCC	TTTGGATGGG	3240
TTACGAACTC	CATCCGGATA	AATGGACGGT	ACAGCCGATC	GTGCTGCCGG	AAAAAGACTC	3300
GTGGACGGTC	AACGACATAC	AGAAGCTCGT	CGGGAAATTG	AATTGGGCGT	CGCAGATTTA	3360

GC AATTATGTAA	ACTCCTTCGC	GGAACGAAAG	CGCTAACGGA	3420
AAG AAGCGGAGCT	CGAACTCGCG	GAAAACCGCG	AGATTCTAAA	3480
TGT ATTACGACCC	GTCGAAAGAC	TTAATCGCGG	AAATACAGAA	3540
GGA CGTATCAAAT	TTATCAAGAG	CCGTTTAAAA	ATCTGAAAAC	3600
TGC GGGGCGCGCA	CACGAACGAC	GTAAAACAAT	TAACGGAGGC	3660
CGG AATCGATCGT	AATATGGGGA	AAGACGCCGA	AATTTAAACT	3720
CGT GGGAAACGTG	GTGGACGGAG	TATTGGCAAG	CGACGTGGAT	3780
TTA ATACGCCGCC	GCTCGTGAAA	TTATGGTACC	AGCTCGAGAA	3840
CGG AAACGTTCTA	CGTCGACGGC	GCGGCGAACC	GCGAGACGAA	3900
ACG TTACGAATCG	CGGACGCCAA	AAAGTCGTCA	CGCTAACGGA	3960
CGG AGTTACAAGC	GATTTATCTC	GCGTTGCAGG	ATTCGGGACT	4020
CGG ACTCGCAATA	CGCGTTAGGA	ATCATTCAAG	CGCAACCGGA	4080
TAG TCAATCAAAT	AATCGAGCAG	TTAATAAAAA	AGGAAAAGGT	4140
CGG CGCACAAAGG	AATCGGCGGA	AACGAACAAG	TCGATAAATT	4200
GGA AAGTACTATT	TTTAGACGGA	ATCGATAAGG	CGCAAGACGA	4260
CGA ATTGGCGCGC	GATGGCGTCG	GATTTTAACC	TGCCGCCGGT	4320
STCG CGTCGTGCG/	A TAAATGTCAG	CTAAAAGGCG	AAGCGATGCA	4380
TCGC CGGGAATATC	GCAACTCGAT	TGTACGCATT	TAGAAGGAAA	4440
STTC ACGTCGCGT	GGGATATATC	GAAGCGGAAG	TTATTCCGGC	4500
ACGG CGTATTTTC	TTTAAAATTA	GCGGGACGAT	GGCCCGTAAA	4560
AATG GCTCGAATT	T CACCGGCGCG	ACGGTTCGCG	CGGCGTGTTG	4620
CAGG AATTCGGAA	T TCCGTACAAT	CCGCAATCGC	AAGGCGTCGT	4680
GAAT TAAAGAAAA	T TATCGGACAG	GTACGCGATC	AGGCGGAACA	4740
CAAA TGGCGGTAT	T CATCCACAAT	TTTAAACGAA	AAGGCGGGAT	4800
GGCG AACGAATCG	T CGACATAATO	GCGACGGACA	TACAAACGAA	4860
ATTA CGAAAATTC	A AAATTTTCG(GTTTATTACO	GCGACTCGCG	4920
GGAC CGGCGAAGC	T CCTCTGGAA	A GGCGAAGGC	G CGGTCGTAAT	4980
ATAA AAGTCGTGC	C GCGACGAAAA	A GCGAAGATC	A TTCGCGATTA	5040
	AAG AAGCGGAGCT TGT ATTACGACCC GGA CGTATCAAAT TGC GGGGCGCGCA CGG AATCGATCGT CGT GGGAAACGTG TTA ATACGCCGCC CGG AAACGTTCTA ACG TTACGAATCG CGG ACTCGCAATA TAG TCAATCAAAT TAG TCAATCAAAT TAG TCAATCAAAT TCGA ATTGGCGCGC GTCG CGCGAATATC TCGA CGTCGTGCGA TCGC CGGGAATATC TCGA CGTCGTGCGA TCGC CGGGAATATC TCGA ATTGGCGCGC TCGC CGGGAATATC TCGA ATTGGCGCGC TCGC CGGGAATATC TCGA ATTCGGAAT TCAATCAAAT TCGA ATTGGCGCGC TCGC CGGGAATATC TCGA ATTCGGAAT TCAATCGAAT TCAATCGAAT TCAATCGAAT TCAAGAAAAT TCAAA TGGCGGTAT TCAAGAAAAT TCAAA TGGCGGTAT TCAAAGAAAAT TCAAA TGGCGGAATCG TCGC CGGCGAAGC TCGC CGGCGAAGC TCGC CGGCGAAGC TCAAAATTCGGAAT TCAAAAATTC TCAAAAAATTC TCAAAAATTC TCAAAAAATTC TCAAAAATTC TCAAAATTC TCAAAATTC TCAAAATTC TCAAAATTC TC	AAG AAGCGGAGCT CGAACTCGCG IGT ATTACGACCC GTCGAAAGAC GGA CGTATCAAAT TTATCAAGAG IGC GGGGCGCGCA CACGAACGAC CGG AATCGATCGT AATATGGGGA CGT GGGAAACGTG GTGGACGGAG TTA ATACGCCGCC GCTCGTGAAA CGG AAACGTTCTA CGTCGACGGC ACG TTACGAATCG CGGACGCCAA CGG ACTCGCAATA CGCGTTAGGA CGG ACTCGCAATA CGCGTTAGGA CGG CGCACAAAGG AATCGGCGGA CGG CGCACAAAGG AATCGGCGGA CGG CGCCCAAAAGG AATCGGCGGA CGG CGCCCCAAAAGG AATCGGCGGA CGG CGCCCCAAAAGG AATCGGCGGA CGG CGCCCCAAAAGG AATCGGCGGA CGG CGTCGTGCGA TAAATGTCAG CTCG CGTCGTGCGA TAAATGTCAG CTCG CGTCGTGCGA TAAATGTCAG CTCG CGTCGTGCGA TAAATGTCAG CTCG CGTCGTGCGA TAAATGTCAG CAGG CGTATTTTCT TTTAAAATTA CATG GCTCGAATTT CACCGGCGCG CAGG AATTCGGAAT TCCGTACAAT CAAA TGGCGGTATT CATCCACAAT CAAA TGGCGGTATT CATCCACAAT CAAA TGGCGGTATT CATCCACAAT CAAA TGGCGGTATT CATCCACAAT CAAA TGGCGGAATCGT CGACATAATC CATTA CGAAAATTCA AAATTTTCGC CAGG CGCGAAACTCA AAATTTTCGC	AGG AAGCGGAGCT CGAACTCGCG GAAAACCGCG IGT ATTACGACCC GTCGAAAGAC TTAATCGCGG IGT ATTACGACCC GTCGAAAGAC TTAATCGCGG IGG CGTATCAAAT TTATCAAGAG CCGTTTAAAA IGC GGGGCGCGCA CACGAACGAC GTAAAACAAT CGG AATCGATCGT AATATGGGGA AAGACGCCGA CGT GGGAAACGTG GTGGACGGAG TATTGGCAAG ITA ATACGCCGCC GCTCGTGAAA TTATGGTACC CGG AAACGTTCTA CGTCGACGGC GCGGCGAACC ACG TTACGAATCG CGGACGCCAA AAAGTCGTCA CGG ACTCGCAATA CGCGTTAGGA ATCATTCAAG ICGG ACTCGCAATA CGCGTTAGGA ATCATTCAAG ICGG ACTCGCAATA CACGGTTAGGA ATCATTCAAG ICGG CGCACAAAAGG AATCGGCGGA AACGAACAAG ICGA AATTGGCGCG GATGGCGTA AACGAACAAG ICGA ATTGGCGCG GATGGCGTA TGTACGAATA ICTC CGTCGTGCGA TAAATGTCAG CTAAAAGGCG ICGC CGGGAATATG GCAACTCGAT TGTACGCATT ICTC ACGTCGCGTC GGGATATATC GAAGCGGAAG ICGG CGTATTTTCT TTTAAAATTA GCGGGACGAT ICAAG GCTCGAATTT CACCGGCGCG ACGGTTCGCG ICAAA TGGCGGAAT TCCGTACAAT CCGCAATCGC ICAAA TGGCGGTATT CACCGGCGCG ACGGTTCGCG ICAAA TGGCGGTATT CACCGGCACG TTAAAACGAA ICAAA TGGCGGTATT CATCCACAAT TTTAAACGAA ICAAA TGGCGGTATT CATCCACAAT TTTAAACGAA ICAAA TGGCGGTATT CATCCACAAT TTTAAACGAA ICAAA TGGCGGAAGCT CGACATAATC GCGACGGACA ICAAA TGGCGGAAGCT CGACATAATC GCGACGGACA ICAAA TGGCGGAAGCT CGACATAATC GCGACGGACAA ICAAA TGGCGGAAGCT CCTCTGGAAA GGCGAAGGCC ICAAC CGGCGAAGCT CCTCTGGAAA GGCGAAGGCC ICAAC CGGCCGAAGCT CCTCTGGAAA GGCGAAGGCC ICAAC CGGCCGAAGCT CCTCTGGAAA GGCGAAGGCC	AAG AAGCGGAGCT CGAACTCGCG GAAAACCGCG AGATTCTAAA TGT ATTACGACCC GTCGAAAGAC TTAATCGCGG AAATACAGAA GGA CGTATCAAAT TTATCAAGAG CCGTTTAAAA ATCTGAAAAC GGA CGTATCAAAT TTATCAAGAG CCGTTTAAAA ATCTGAAAAC GGA CGTATCAAAT TTATCAAGAG CCGTTTAAAA ATCTGAAAAC GGA CGGGCGCGCA CACGAACGAC GTAAAACAAT TAACGGAGGC CGG AATCGATCGT AATATGGGGA AAGACGCCGA AATTTAAACT CGT GGGAAACGTG GTGGACGGAG TATTGGCAAG CGACGTGGAT TTA ATACGCCGCC GCTCGTGAAA TTATGGTACC AGCTCGAGAA CGG AAACGTTCTA CGTCGACGGC GCGGCGAACC GCGAGACGAA ACG TTACGAATCG CGGACGCCAA AAAGTCGTCA CGCTAACGGA CGG ACTCGCAATA CGCGTTAGGA ATCATTCAAG CGCAACCGGA TAG TCAATCAAAT AATCGAGCAG TTAATAAAAA AGGAAAAGGT CGG CGCACAAAGG AATCGGCGGA AACGAACAAG TCGATAAATT CGGA AAGTACTATT TTTAGACGGA ATCGATAAGG CGCAAGACGA CGGA ATTGGCGCGC GATGGCGTCG GATTTTACC TGCCGCCGGT CTCG CGTCGTGCGA TAAATGTCAG CTAAAAGGCG AAGCGATGCA CTCG CGGGAATATG GCAACTCGAT TGTACGCATT TAGAAGGAAA CTCC CGGGGAATTTC CACCGGCCGG ACGGTTCGCG CGGCCGTAAA CTCG CGGGAATTTC TTTAAAATTA GCGGGACGAT GGCCCGTAAA CTCG CGGGGAATTT CACCGGCCGG ACGGTTCGCG CGGCGTGTTG CAGG CATTTTTCT TTTAAAATTA GCGGAACGAT GGCCCGTAAA CTCAA TTAGAAGAAAT TCCGTACAAT CCGCAATCGC AAGGCGGACA CAAA TGGCGGTATT CATCCACAAT TTTAAACGAA AAGGCGGGAT CAAA TGGCGGTATT CATCCACAAT TTTAAACGAA AAGGCGGGAT CAAA TGGCGGTATT CATCCACAAT TTTAAACGAA AAGGCGGGAT CAAAA TGGCGGAACTC CGGCACGAAAA GCGAAGACCG CAAAAATTCA AAATTTCGG GTTTATTACC GCGACTCGCG GGAC CGGCGAAGCT CCTCTGGAAA GCGAAGACCA TACAAACGAA ATTA CGAAAATTCA AAATTTTCGC GTTTATTACC GCGACTCGCG GGAC CGGCGAAGCT CCTCTGGAAAA GCGAAGACCA TTCCACACTT ATAA AAGTCGTGCC CGCGCGAAAAA GCGAAGACCA TTCCACATT

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TGGAAAACAG	ATGGCAGGTG	ATGATTGTGT	GGCAAGTAGA	CAGGATGAGG	ATTCGCACGT	5100
GGAAATCGCT	CGTAAAACAC	CATATGTACG	TTTCGGGGAA	AGCGCGCGGA	TGGTTTTATC	5160
GCCATCACTA	CGAATCGCCG	CATCCGCGCA	TATCGTCGGA	AGTACACATC	CCGCTCGGGG	5220
ATGCGCGCCT	CGTAATAACG	ACGTATTGGG	GTCTGCATAC	GGGCGAACGC	GACTGGCATC	5280
TCGGTCAGGG	CGTCTCGATC	GAATGGCGCA	AAAAGCGCTA	TTCGACGCAA	GTCGACCCGG	5340
AACTCGCGGA	CCAACTAATT	CATCTGTATT	ACTTCGACTG	TTTTTCGGAC	TCGGCGATAC	5400
GCAAGGCGTT	ACTCGGACAC	ATCGTTTCGC	CGCGCTGCGA	ATATCAAGCG	GGACATAACA	5460
AGGTCGGATC	GCTACAATAC	CTCGCGCTCG	CGGCGTTAAT	AACGCCGAAA	AAGATAAAGC	5520
CGCCGTTGCC	GTCGGTTACG	AAACTGACGG	AGGATCGATG	GAACAAGCCC	CAGAAGACCA ·	5580
AGGGCCACAG	AGGGAGCCAC	ACAATGAATG	GACACTAGAG	CTTTTAGAGG	AGCTTAAGAA	5640
CGAAGCGGTT	CGCCATTTTC	CGCGCATTTG	GCTCCACGGC	TTAGGGCAAC	ATATCTACGA	5700
AACGTATGGG	GATACGTGGG	CGGGCGTCGA	AGCGATAATA	AGAATTCTGC	AACAACTGCT	5760
GTTTATCCAT	TTTCAGAATT	GGGTGTCGAC	GTAGCAGAAT	AGGCGTTACT	CGACAGAGGA	5820
GAGCAAGAAA	TGGAGCCAGT	AGATCCTAGA	CTAGAGCCCT	GGAAGCATCC	AGGAAGTCAG	5880
CCTAAAACTG	CTTGTACCAA	TTGCTATTGT	AAAAAGTGTT	GCTTTCATTG	CCAAGTTTGT	5940
TTCATAACAA	AAGCCTTAGG	CATCTCCTAT	GGCAGGAAGA	AGCGGAGACA	GCGACGAAGA	6000
GCTCATCAGA	ACAGTCAGAC	TCATCAAGCT	TCTCTATCAA	AGCAGTAAGT	AGTACATGTA	6060
ACGCAACCTA	TACCAATAGT	AGCAATAGTA	GCATTAGTAG	TAGCAATAAT	AATAGCAATA	6120
GTTGTGTGGT	CCATAGTAAT	CATAGAATAT	AGGAAAATAT	TAAGACAAAG	AAAAATAGAC	6180
AGGTTAATTG	ATAGACTAAT	AGAAAGAGCA	GAAGACAGTG	GCAATGCGAG	TGAAGGAGAA	6240
ATATCAGCAC	TTGTGGCGAT	GGGGGTGGCG	ATGGGGCACG	ATGCTCCTCG	GGATGTTGAT	6300
GATCTGTTCG	GCTACGGAAA	AATTGTGGGT	CACGGTCTAT	TACGGCGTAC	CGGTGTGGAA	6360
GGAAGCGACG	ACGACGCTAT	TTTGCGCGTC	GGACGCGAAA	GCGTACGATA	CGGAGGTACA	6420
TAACGTTTGG	GCGACGCATG	CGTGCGTACC	GACGGACCCG	AACCCGCAAG	AAGTCGTACT	6480
CGTAAACGTG	ACGGAAAATT	TCGACATGTG	GAAAAACGAC	ATGGTCGAAC	AGATGCATGA	6540
GGATATAATC	TCGTTATGGG	ATCAATCGCT	AAAGCCGTGC	GTAAAATTAA	CGCCGCTCTG	6600
CGTTTCGTTA	AAGTGCACGG	ATTTGAAGAA	TGATACGAAT	ACGAATTCGT	CGTCGGGGCG	6660
AATGATAATG	GAGAAAGGCG	AGATAAAAAA	CTGCTCGTTC	AATATCTCGA	CGTCGATACG	6720

(CGGTAAGGTG	CAGAAAGAAT	ACGCGTTTTT	TTATAAACTE	GATATAATAC	CGATCGATAA	6780
(CGATACGACG	TCGTATTCGT	TGACGTCGTG	TAACACGTCG	GTCATTACGC	AGGCGTGTCC	6840
(SAAGGTATCG	TTCGAGCCGA	TTCCGATACA	TTATTGTGCG	CCGGCGGGTT	TCGCGATTCT	6900
1	AAAATGTAAT	AATAAGACGT	TCAACGGAAC	GGGACCGTGT	ACGAACGTCT	CGACGGTACA	6960
1	ATGTACGCAC	GGAATTCGGC	CGGTCGTATC	GACGCAACTG	CTGTTAAACG	GCTCGCTCGC	7020
(GGAAGAAGAG	GTCGTAATTC	GATCGGTCAA	TTTCACGGAC	AACGCGAAAA	CGATAATCGT	7080
,	ACAGCTGAAC	ACGTCGGTCG	AAATTAATTG	TACGCGACCG	AACAACAATA	CGCGAAAACG	7140
,	AATCCGTATC	CAGCGCGGAC	CGGGGCGCGC	ATTCGTTACG	ATCGGAAAAA	TCGGAAATAT	7200
	GCGACAAGCG	CATTGTAACA	TTTCGCGCGC	GAAATGGAAT	AACACGTTAA	AACAGATCGA	7260
	TTCGAAATTA	CGCGAACAAT	TCGGAAATAA	TAAAACGATA	ATCTTTAAGC	AATCGTCGGG	7320
	CGGCGACCCG	GAAATCGTAA	CGCACTCGTT	TAATTGTGGC	GGCGAATTIT	TCTACTGTAA	7380
	TTCGACGCAA	CTGTTTAATT	CGACGTGGTT	TAATTCGACG	TGGTCGACGG	AAGGGTCGAA	7440
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	GTGGCAGAAA	GTCGGAAAAG	CGATGTACGC	GCCGCCGATC	TCGGGACAAA	TTCGATGTTC	7560
	GTCGAATATT	ACGGGGCTGC	TATTAACGCG	CGACGGCGGT	AATTCGAACA	ACGAGTCCGA	7620
•	GATCTTCCGA	CTCGGCGGCG	GCGATATGCG	CGACAATTGG	CGATCGGAAT	ATAAATA	7680
	TAAAGTCGTA	AAAATCGAAC	CGCTCGGCGT	CGCGCCGACG	AAGGCGAAGC	GACGCGTCGT	7740
	GCAGCGCGAA	AAACGCGCGG	TCGGAATCGG	CGCGTTGTTC	CTCGGGTTCC	TCGGCGCGGC	7800
	CGGATCGACG	ATGGGCGCGG	CGTCGATGAC	GCTGACGGTA	CAGGCGCGAC	AATTATTGTC	7860
	GGGTATCGTG	CAGCAGCAGA	ACAATTTGCT	GCGCGCTATC	GAGGCGCAAC	AGCATCTGTT	7920
	GCAACTCACG	GTCTGGGGCA	TCAAGCAGCT	CCAAGCGCGA	ATCCTCGCGG	TCGAACGATA	7980
	CCTAAAGGAT	CAACAGCTCC	TCGGGATTTG	GGGTTGCTCG	GGAAAACTCA	TTTGCACGAC	8040
	GGCGGTGCCG	TGGAATGCGT	CGTGGTCGAA	TAAATCGCTC	GAACAGATCT	GGAATCACAC	8100
	GACGTGGATG	GAGTGGGACC	GCGAAATTAA	CAATTACACG	TCGTTAATAC	ACTCGTTAAT	8160
	TGAAGAATCG	CAAAACCAGO	: AAGAAAAGA	A TGAACAAGAA	TTACTCGAAC	C TCGATAAATG	8220
						A AATTATTCAT	8280
	AATGATCGTC	GGCGGCCTCG	TCGGTTTACC	AATCGTTTT(GCGGTACTT	CGATCGTGAA	8340
	TCGCGTTCGC	G CAGGGATAT	CGCCGTTATO	GTTTCAGACO	CACCTCCCA	A TCCCGAGGGG	8400

ACCCGACAGG	CCCGAAGGAA	TAGAAGAAGA	AGGTGGAGAG	AGAGACAGAG	ACAGATCCAT	8460
TCGATTAGTG	AACGGATCCT	TGGCACTTAT	CTGGGACGAT	CTGCGGAGCC	TGTGCCTCTT	8520
CAGCTACCAC	CGCTTGAGAG	ACTTACTCTT	GATTGTAACG	AGGATTGTGG	AACTTCTGGG	8580
ACGCAGGGGG	TGGGAAGCCC	TCAAATATTG	GTGGAATCTC	CTACAGTATT	GGAGTCAGGA	8640
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AGATAGGGTT	ATAGAAGTAG	TACAAGGAGC	TTGTAGAGCT	ATTCGCCACA	TACCTAGAAG	8760
AATAAGACAG	GGCTTGGAAA	GGATTTTGCT	ATAAGATGGG	CGGCAAGTGG	TCGAAATCGT	8820
CGGTGATTGG	ATGGCTTACG	GTACGCGAAC	GCATGCGCCG	CGCCGAGCCG	GCGGCGGACG	8880
GCGTCGGCGC	CGCGTCGCGC	GACCTGGAAA	AACACGGCGC	GATCACGTCG	TCGAACACGG	8940
CGGCGACGAA	CGCGGCGTGC	GCGTGGCTCG	AAGCGCAAGA	GGAGGAGGAG	GTCGGTTTTC	9000
CGGTCACGCC	GCAGGTACCG	TTACGCCCGA	TGACGTACAA	GGCGGCGGTC	GATCTTTCGC	9060
ACTTTTTAAA	AGAAAAGGGC	GGACTCGAAG	GGCTAATTCA	CTCGCAACGC	CGCCAAGATA	9120
TCCTCGATCT	GTGGATCTAC	CACACGCAAG	GCTACTTCCC	GGATTGACAG	AACTACACAC	9180
CAGGGCCAGG	GGTCAGATAT	CCACTGACCT	TTGGATGGTG	CTACAAGCTA	GTACCAGTTG	9240
AGCCAGÁTAA	GATAGAAGAG	GCCAATAAAG	GAGAGAACAC	CAGCTTGTTA	CACCCTGTGA	9300
GCCTGCATGG	GATGGATGAC	CCGGAGAGAG	AAGTGTTAGA	GTGGAGGTTT	GACAGCCGCC	9360
TAGCATTTCA	TCACGTGGCC	CGAGAGCTGC	ATCCGGAGTA	CTTCAAGAAC	TGCTGACATC	9420
GAGCTTGCTA	CAAGGGACTT	TCCGCTGGGG	ACTTTCCAGG	GAGGCGTGGC	CTGGGCGGGA	9480
CTGGGGAGTG	GCGAGCCCTC	AGATCCTGCA	TATAAGCAGC	TGCTTTTTGC	CTGTACTGGG	9540
TCTCTCTGGT	TAGACCAGAT	CTGAGCCTGG	GAGCTCTCTG	GCTAACTAGG	GAACCCACTG	9600
CTTAAGCCTC	AATAAAGCTT	GCCTTGAGTG	CTTCAAGTAG	TGTGTGCCCG	TCTGTTGTGT	9660
GACTCTGGTA	ACTAGAGATO	CCTCAGACCC	TTTTAGTCAG	TGTGGAAAAT	CTCTAGCA	9718

THAT WHICH IS CLAIMED IS:

- 1. An attenuated virus, not naturally occurring, containing at least 1 additional methylation site in the genome of said virus compared to the corresponding wild-type virus.
- 2. An attenuated virus according to claim 1, said virus comprising a viral capsid containing said genome.
- 3. An attenuated virus of claim 1, containing at least 10 additional methylation sites over the 10 corresponding wild-type virus.
 - 4. An attenuated virus of claim 1, containing at least 100 additional methylation sites over the corresponding wild-type virus.
- 5. An attenuated virus of claim 1 wherein said
 15 methylation site is a CG segment.
 - 6. An attenuated virus according to claim 1, wherein said virus is a DNA virus.
 - 7. An attenuated virus according to claim 1, wherein said virus is a retrovirus.
- 8. An attenuated virus of claim 1 wherein said virus is a retrovirus selected from the group consisting of B-type retroviruses, C-type retroviruses, D-type retroviruses, Lentiviruses, T-cell leukemia viruses, and foamy viruses.
- 9. An attenuated virus of claim 1, wherein said virus is HIV-1.

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10. An attenuated virus of claim 1, wherein said virus is SIV.

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- 11. An attenuated virus of claim 1, wherein said virus is HTLV-1.
- 5 12. An attenuated virus of claim 1, wherein said virus is a retrovirus and wherein an attenuating deletion mutation is included therein.
 - 13. A DNA encoding a virus of claim 1.
- 14. An expression vector containing a DNA of 10 claim 13.
 - 15. An expression vector of claim 14, wherein said expression vector is a Baculovirus.
- 16. A host cell containing a DNA of claim 13 and capable of expressing the encoded virus, which host cell does not methylate said DNA sufficient to inactivate the expression of the encoded viral genome.
 - 17. A host cell according to claim 16, which host cell lacks capacity to methylate DNA because of treatment of said host cell with a methylation inhibitor.
- 20 18. A host cell according to claim 17 wherein said methylation inhibitor is 5-azadeoxycytidine or 5-azacytidine.
- 19. A pharmaceutical formulation comprising a virus according to claim 1 in combination with a pharmaceutically acceptable carrier.
 - 20. A formulation according to claim 19, wherein said formulation is an oral formulation.

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- A formulation according to claim 19, 21. wherein said formulation is a parenterally injectable vaccine formulation.
- A formulation according to claim 19, 22. 5 wherein said formulation is an inhalation formulation.
 - A method of producing an immune response in a subject, comprised of administering a virus of claim 1 to said subject in an amount effective to produce an immune response in said subject.
- A method according to claim 23, wherein 24. 10 said administering step is carried out by orally administering said virus to said subject.
- A method according to claim 23, wherein said administering step is carried out by parenterally 15 injecting said virus into said subject.
 - A method according to claim 23, wherein said subject is an animal subject.
 - 27. A method according to claim 23, wherein said subject is a human subject.
- 28. A method of making an attenuated virus, 20 not naturally occurring, containing at least 1 additional methylation site in the genome of said virus compared to corresponding wild-type virus; said method the comprising:
- providing a host cell containing an expression 25 vector, said expression vector containing a DNA encoding said attenuated virus, which host cell does not methylate said DNA sufficient to inactivate the expression of the encoded viral genome; and

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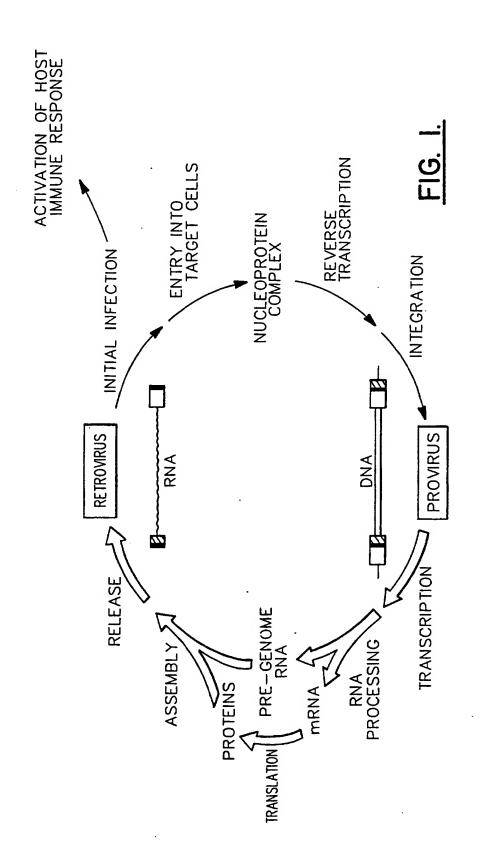
expressing said attenuated virus in said host cell.

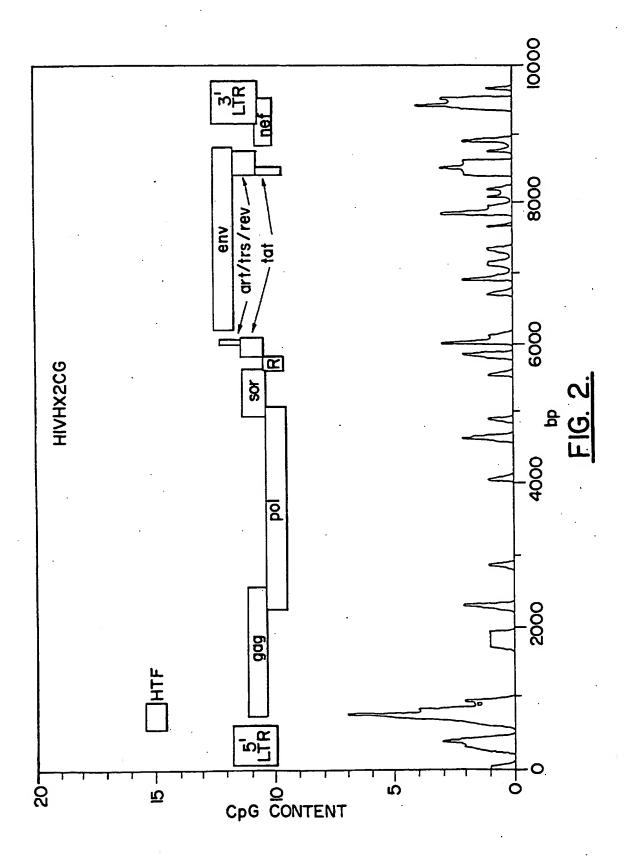
- 29. A method according to claim 28, the genome of said virus containing at least 10 additional methylation sites over the corresponding wild-type virus.
 - 30. A method according to claim 28, wherein said virus is a DNA virus.
 - 31. A method according to claim 28, wherein said virus is a retrovirus.
- 32. A method according to claim 28, wherein said expression vector is a Baculovirus.
 - 33. A method according to claim 28, wherein said host cell is an insect cell.
- 34. A method according to claim 28, wherein 15 said host cell is a mammalian cell.
 - 35. An oligonucleotide probe useful for distinguishing between (i) an attenuated virus, not naturally occurring, containing at least 1 additional methylation site in the genome of said virus compared to the corresponding wild-type virus, and (ii) said corresponding wild-type virus, said oligonucleotide probe selected from the group consisting of:
- (a) oligonucleotide probes that selectively hybridize to the nucleic acid of an attenuated virus of
 (i) above, and which do not hybridize to the nucleic acid of the wild-type virus of (ii) above under the same hybridization conditions; and
- (b) oligonucleotide probes that selectively hybridize to the nucleic acid of a wild-type virus of 30 (ii) above, and which do not hybridize to the nucleic

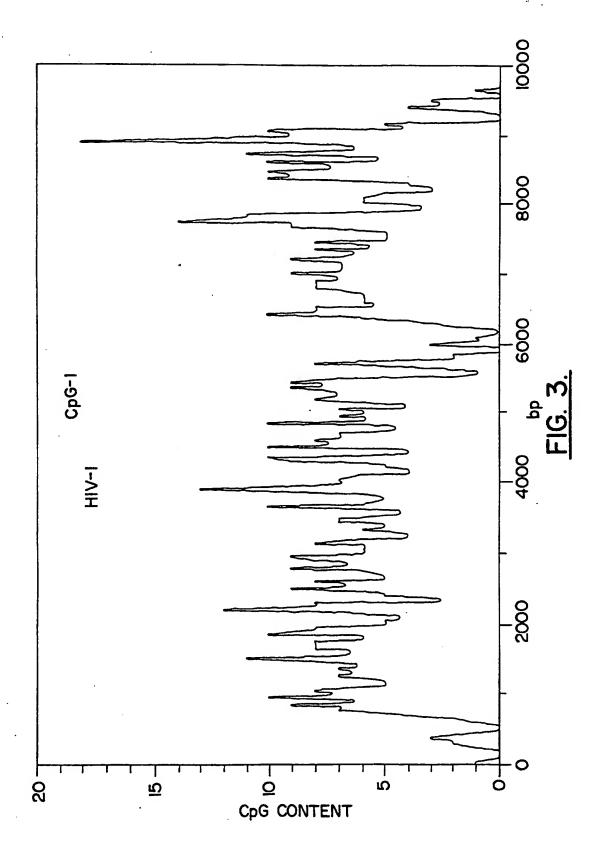
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acid of the attenuated virus of (i) above under the same hybridization conditions.

- 36. An oligonucleotide probe according to claim 35 conjugated to a detectable group.
- 5 37. An oligonucleotide probe according to claim 35, wherein said probe is a PCR extension primer.







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IPC 6	C12N5/10	76 C12N7/04	C12Q1/70
·	o International Patent Classification (IPC) or to both national class	ification and IPC	
	SEARCHED ocumentation searched (classification system followed by classifica-	ton nembols)	
IPC 6	C12N A61K C12Q		
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Electronic d	iata base consulted during the international search (name of data ba	se and, where practical, search terms	s used)
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	an the priority date claimed	"&" document member of the same	patent family
Date of the	actual completion of the international search	Date of mailing of the internati	·
8	February 1996	15. a	3. 96
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